1600





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RAW SEQUENCE LISTING

DATE: 01/31/2002

PATENT APPLICATION: US/10/033,026

TIME: 10:29:08

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        Schorge, Stephanie
 6 <120> TITLE OF INVENTION: HUMAN N-TYPE CALCIUM CHANNEL ISOFORM AND USES THEREOF
 8 <130> FILE REFERENCE: B1055/7000
10 <140> CURRENT APPLICATION NUMBER: 10/033,026
11 <141> CURRENT FILING DATE: 2001-12-28
13 <150> PRIOR APPLICATION NUMBER: US 09/268,163
14 <151> PRIOR FILING DATE: 1999-03-12
16 <150> PRIOR APPLICATION NUMBER: US 60/077,901
17 <151> PRIOR FILING DATE: 1998-03-13
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52 gggatgcacg cggggcccgg gagcc atg gtc cgc ttc ggg gac gag ctg ggc
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56 Gly Arg Tyr Gly Gly Pro Gly Gly Gly Glu Arg Ala Arg Gly Gly Gly
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268
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PATENT APPLICATION: US/10/033,026

DATE: 01/31/2002 TIME: 10:29:08

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66	Deu	1 7 1	60	110	110	110	,	65	01		010		70			5	
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68					Phe												
69	Ser	75	rne	Vai	rne	DCI	80	p	11011	,	, 41	85	_10	-1-		-1-	
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7 <i>7</i>	ASP	пуъ	1111	125	Mec	per	GIU	пта	130	пор	nsp	1111	014	135	-1-	1110	
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80	TIO	211	Tlo	Dho	Cys	Dha	Glu	λla	999 999	Tle	Lvs	Tle	Tle	Ala	Len	Glv	001
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	Phe	155	PILE	птэ	пуз	Gry.	160	1 Y 1	пец	nrg	non	165		11011	,	1100	
84	~~~		~+ ~	a+ a	gtc	ata		aaa	ato	ott	acc		act	απа	act	gac	700
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91	a+ a	a+a	+ a+	aaa	att	002	a art	t+a	aaa		αtα	ctc	aan	tcc		atσ	796
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97	2+0	ata		+++	gcc	ato	att		cta	σασ	ttc	tac		aac	ааσ	ttc	892
98					Ala												07-
99	TIE	235	Mec	FIIC	AIu	116	240	GLY	пси	Olu	1110	245	1100		_,_	10	
100	C D I		a ac	r tai	- ++/			- aσ	r aca	a σai	t ac		T CC	a ata	ı aat	t gac	940
101	Ui	e Tare	9 9C	a Cv	Dhe	Dr	) A G I	n Sei	r Thi	r Ası	n Ala	a Gli	ı Pro	o Vai	Gl	y Asp	
101	250		o vic	ı cy.	5 1 110	25					260					265	
102			a tai	- ~~				י ככי	a aca	י כמו			r σac	r aa	r σac	act	988
103																Thr	
104	F110	= PI	o cy:	s Gr	у шуг 270		~ VJT(		~ 13±6	27					280		
105	(r = 1	7 + A	0 000	יביה ד		-	T CC:	a aa:	3 000			ר ממי	ata	. acc		ttt	1036
107	ga!	9 CT	e Are	יונים די	<sub>1</sub> Γαι	- U	) Dr	- ሃሃ ጉርጉ	r Pro	n Aes	n Dh	יום ב	, T1	- ∽ov	- Asi	n Phe	2020
107	GI	и су:	2 WT	28!		- 11j	, ,,	. GI	290			- J <sub>-</sub>	, ++	295	5		
T00				20.	,				2)(	•				٠,٠	-		

PATENT APPLICATION: US/10/033,026

DATE: 01/31/2002 TIME: 10:29:08

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133 134 135	Asn	gac Asp	Leu	Ile	His 430	Ala	Glu	Glu	Gly	Glu 435	Asp	Arg	Phe	Ala	Asp 440	Leu	1468
136 137 138		gct Ala															1516
139 140 141		gag Glu															1564
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145 146 147		gtg Val															1660
148 149 150	cag Gln	ccg Pro	cgg Arg	cgg Arg	ctt Leu 510	acc Thr	acg Thr	acc Thr	ctg Leu	tat Tyr 515	ttt Phe	gca Ala	gag Glu	ttt Phe	gtt Val 520	ttc Phe	1708
151 152 153		ggt Gly															1756
154 155 156		aga Arg		tac													1804
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PATENT APPLICATION: US/10/033,026

DATE: 01/31/2002 TIME: 10:29:08

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161       Ser       Ser       Phe       Gly       Ile       Ser       Val       Leu       Arg       Ala       Leu       Arg       Leu       Arg       Leu       Arg       Leu       Arg       Leu       Arg       Ile       Arg       Ile       Arg       Ile       Ile       Arg       Ile       I	1948 1996
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ttc aaa gtc acg aag tac tgg agc tcc ctg cgg aac ctg gtg gtg tcc  164 Phe Lys Val Thr Lys Tyr Trp Ser Ser Leu Arg Asn Leu Val Val Ser  165 590 595 600  166 ctg ctg aac tcc atg aag tcc atc atc agc ctg ctc ttc ttg ctc ttc  167 Leu Leu Asn Ser Met Lys Ser Ile Ile Ser Leu Leu Phe Leu Leu Phe  168 605 610 615  169 ctg ttc att gtg gtc ttc gcc ctg ctg ggg atg cag ctg ttt ggg gga  170 Leu Phe Ile Val Val Phe Ala Leu Leu Gly Met Gln Leu Phe Gly Gly	1996
164       Phe Lys Val Thr Lys Tyr Trp Ser Ser Leu Arg Asn Leu Val Val Ser         165       590       595       600         166       ctg ctg aac tcc atg aag tcc atc atc agc ctg ctc ttc ttg ctc ttc         167       Leu Leu Asn Ser Met Lys Ser Ile Ile Ser Leu Leu Phe Leu Leu Phe         168       605       610         609       615         169       ctg ttc att gtg gtc ttc gcc ctg ctg ggg atg cag ctg ttt ggg gga         170       Leu Phe Ile Val Val Phe Ala Leu Leu Gly Met Gln Leu Phe Gly Gly	1996
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2/2	2092
172 cag ttc aac ttc cag gat gag act ccc aca acc aac ttc gac acc ttc	2092
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182 Gly Met Phe Ser Ser Phe Tyr Phe Ile Val Leu Thr Leu Phe Gly Asn	
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184 tac act ctg ctg aat gtc ttt ctg gcc atc gct gtg gac aac ctg gcc	2284
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185 Tyr Thr Leu Leu Asn Val Phe Leu Ala Ile Ala Val Asp Asn Leu Ala	
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188 Asn Ala Gln Glu Leu Thr Lys Asp Glu Glu Glu Met Glu Glu Ala Ala	
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189 715 720 725 190 aat cag aag ctt gct ctg caa aag gcc aaa gaa gtg gct gaa gtc agc	2380
	2380
190 aat cag aag ctt gct ctg caa aag gcc aaa gaa gtg gct gaa gtc agc	2380
190 aat cag aag ctt gct ctg caa aag gcc aaa gaa gtg gct gaa gtc agc 191 Asn Gln Lys Leu Ala Leu Gln Lys Ala Lys Glu Val Ala Glu Val Ser 192 730 745	2380 2428
190 aat cag aag ctt gct ctg caa aag gcc aaa gaa gtg gct gaa gtc agc 191 Asn Gln Lys Leu Ala Leu Gln Lys Ala Lys Glu Val Ala Glu Val Ser 192 730 735 740 745 193 ccc atg tct gcc gcg aac atc tcc atc gcc gcc agg cag cag aac tcg	
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190 aat cag aag ctt gct ctg caa aag gcc aaa gaa gtg gct gaa gtc agc 191 Asn Gln Lys Leu Ala Leu Gln Lys Ala Lys Glu Val Ala Glu Val Ser 192 730 735 740 745 193 ccc atg tct gcc gcg aac atc tcc atc gcc gcc agg cag cag aac tcg 194 Pro Met Ser Ala Ala Asn Ile Ser Ile Ala Ala Arg Gln Gln Asn Ser 195 750 755 760 196 gcc aag gcg cgc tcg gtg tgg gag cag cgg gcc agc cag cta cgg ctg	
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PATENT APPLICATION: US/10/033,026

DATE: 01/31/2002 TIME: 10:29:08

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210
212       Pro Glu Gly Val Asp Pro Pro Arg Arg His His Arg His Arg Asp Lys         213       845       850       855         214       gac aag acc ccc gcg gcg ggg gac cag gac cag gac cga gag gcc ccg aag       2764         215       Asp Lys Thr Pro Ala Ala Gly Asp Gln Asp Arg Ala Glu Ala Pro Lys       860       865       870         217       gcg gag agc ggg gag ccc ggt gcc cgg gag gag cgg ccg ccg ccg cac       2812         218       Ala Glu Ser Gly Glu Pro Gly Ala Arg Glu Glu Arg Pro Arg Pro His       880       885         220       cgc agc cac agc aag gag gcc gcg ggg gcc ccg ggg gg
213
214 gac aag acc ccc gcg gcg ggg gac cag gac cga gca gag gcc ccg aag 2764 215 Asp Lys Thr Pro Ala Ala Gly Asp Gln Asp Arg Ala Glu Ala Pro Lys 216 860 865 870 217 gcg gag agc ggg gag ccc ggt gcc cgg gag gag cgg ccg cgg ccg cac 2812 218 Ala Glu Ser Gly Glu Pro Gly Ala Arg Glu Glu Arg Pro Arg Pro His 219 875 880 885 220 cgc agc cac agc aag gag gcc gcg ggg ccc ccg gag gcg cgg agc gag 2860 221 Arg Ser His Ser Lys Glu Ala Ala Gly Pro Pro Glu Ala Arg Ser Glu 222 890 895 900 905 223 cgc ggc cga ggc cca ggc ccc gag ggc ggc
215       Asp Lys Thr Pro Ala Ala Gly Asp Gln Asp Arg Ala Glu Ala Pro Lys         216       860       865       870         217       gcg gag agc ggg gag ccc ggt gcc cgg gag gag cgg ccg ccg ccg ccac       2812         218       Ala Glu Ser Gly Glu Pro Gly Ala Arg Glu Glu Arg Pro Arg Pro His       885         219       875       880       885         220       cgc agc cac agc aag gag gcc gcg ggg ccc ccg gag gcg cgg agc gag       2860         221       Arg Ser His Ser Lys Glu Ala Ala Gly Pro Pro Glu Ala Arg Ser Glu         222       890       895       900       905         223       cgc ggc cga ggc cca ggc cca ggc ccc gag ggc ggc
216       860       865       870         217       gcg gag agc ggg gag ccc ggt gcc cgg gag gag ccg ccg ccg cac       2812         218       Ala Glu Ser Gly Glu Pro Gly Ala Arg Glu Glu Arg Pro Arg Pro His       2812         219       875       880       885         220       cgc agc cac agc aag gag gcc gcg ggg ccc ccg gag gcg cgg agc gag gag
216       860       865       870         217       gcg gag agc ggg gag ccc ggt gcc cgg gag gag ccg ccg ccg cac       2812         218       Ala Glu Ser Gly Glu Pro Gly Ala Arg Glu Glu Arg Pro Arg Pro His       2812         219       875       880       885         220       cgc agc cac agc aag gag gcc gcg ggg ccc ccg gag gcg cgg agc gag gag
218 Ala Glu Ser Gly Glu Pro Gly Ala Arg Glu Glu Arg Pro Arg Pro His 219 875 880 885  220 cgc agc cac agc aag gag gcc gcg ggg ccc ccg gag gcg cgg agc gag 2860 221 Arg Ser His Ser Lys Glu Ala Ala Gly Pro Pro Glu Ala Arg Ser Glu 222 890 895 900 905  223 cgc ggc cga ggc cca ggc ccc gag ggc ggc
219 875 880 885  220 cgc agc cac agc aag gag gcc gcg ggg ccc ccg gag gcg cgg agc gag 2860  221 Arg Ser His Ser Lys Glu Ala Ala Gly Pro Pro Glu Ala Arg Ser Glu  222 890 895 900 905  223 cgc ggc cga ggc cca ggc ccc gag ggc ggc
220 cgc agc cac agc aag gag gcc gcg ggg ccc ccg gag gcg cgg agc gag 221 Arg Ser His Ser Lys Glu Ala Ala Gly Pro Pro Glu Ala Arg Ser Glu 222 890 895 900 905 223 cgc ggc cga ggc cca ggc ccc gag ggc ggc
221 Arg Ser His Ser Lys Glu Ala Ala Gly Pro Pro Glu Ala Arg Ser Glu 222 890 895 900 905  223 cgc ggc cga ggc cca ggc ccc gag ggc ggc
222 890 895 900 905  223 cgc ggc cga ggc cca ggc ccc gag ggc ggc
223 cgc ggc cga ggc cca ggc ccc gag ggc ggc
224 Arg Gly Arg Gly Pro Gly Pro Glu Gly Gly Arg Arg His His Arg Arg 225 910 915 920 226 ggc tcc ccg gag gag gcc gag cgc gag ccc cga cgc cac cgc gcg 227 Gly Ser Pro Glu Glu Ala Ala Glu Arg Glu Pro Arg Arg His Arg Ala 228 925 930 935
225       910       915       920         226       ggc tcc ccg gag gag gcg gcc gag cgg gag ccc cga cgc cac cgc gcg       2956         227       Gly Ser Pro Glu Glu Ala Ala Glu Arg Glu Pro Arg Arg His Arg Ala       228         925       930       935
226 ggc tcc ccg gag gag gcc gag cgg gag ccc cga cgc cac cgc gcg 2956 227 Gly Ser Pro Glu Glu Ala Ala Glu Arg Glu Pro Arg Arg His Arg Ala 228 925 930 935
227 Gly Ser Pro Glu Glu Ala Ala Glu Arg Glu Pro Arg Arg His Arg Ala 228 925 930 935
227 Gly Ser Pro Glu Glu Ala Ala Glu Arg Glu Pro Arg Arg His Arg Ala 228 925 930 935
220
229 cac cgg cac cag gat ccg agc aag gag tgc gcc ggc gcc aag ggc gag 3004
230 His Arg His Gln Asp Pro Ser Lys Glu Cys Ala Gly Ala Lys Gly Glu
231 940 945 950
232 cgg cgc gcg cac cgc ggc ggc ccc cga gcg ggg ccc cgg gag gcg 3052
233 Arg Arg Ala Arg His Arg Gly Gly Pro Arg Ala Gly Pro Arg Glu Ala
234 955 960 965
235 gag agc ggg gag gag ccg gcg cgg cac cgg gcc cgg cac aag gcg 3100
236 Glu Ser Gly Glu Glu Pro Ala Arg Arg His Arg Ala Arg His Lys Ala
237 970 975 980 985
238 cag cot got cac gag got gtg gag aag gag acc acg gag aag gag goc 3148
239 Gln Pro Ala His Glu Ala Val Glu Lys Glu Thr Thr Glu Lys Glu Ala
240 990 995 1000
241 acg gag aag gag gct gag ata gtg gaa gcc gac aag gaa aag gag ctc 3196
242 Thr Glu Lys Glu Ala Glu Ile Val Glu Ala Asp Lys Glu Lys Glu Leu
243 1005 1010 1015
244 cgg aac cac cag ccc cgg gag cca cac tgt gac ctg gag acc agt ggg 3244
245 Arg Asn His Gln Pro Arg Glu Pro His Cys Asp Leu Glu Thr Ser Gly
246 1020 1025 1030
247 act gtg act gtg ggt ccc atg cac aca ctg ccc agc acc tgt ctc cag 3292
248 Thr Val Thr Val Gly Pro Met His Thr Leu Pro Ser Thr Cys Leu Gln
249 1035 1040 1045
250 aag gtg gag gaa cag cca gag gat gca gac aat cag cgg aac gtc act 3340
251 Lys Val Glu Glu Gln Pro Glu Asp Ala Asp Asn Gln Arg Asn Val Thr
252 1050 1055 1060 1065
253 cgc atg ggc agt cag ccc cca gac ccg aac act att gta cat atc cca 3388
254 Arg Met Gly Ser Gln Pro Pro Asp Pro Asn Thr Ile Val His Ile Pro
255 1070 1075 1080

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/033,026

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